

Complex I Human Protein Quantity Dipstick Assay Kit

MS131

Rev.2

DESCRIPTION

Complex I Human Protein Quantity Dipstick Assay Kit

Sufficient materials for 30 (MS131-30) or 90 (MS131-90) measurements.

Kit Contents:

Item	MS131-30	MS131-90
Dipsticks	30	90
Gold-conjugated antibody (dried in microplate wells)	30 wells	90 wells
Buffer A* (Extraction buffer)	15 mL	45 mL
Buffer B* (Blocking buffer)	1 mL	3 mL
Buffer C* (Wash buffer)	1 mL	3 mL

Storage:

Store dipsticks and microplate with gold-conjugated antibody at room temperature out of direct sunlight in their provided containers. High humidity conditions should be avoided. Store Buffers A, B, and C at 4°C, or at -20°C for long-term storage. **Buffers A, B, and C are interchangeable with Buffers A, B, and C from other Quantity Dipstick Assay Kits (MS133, MS431, MSF31, MSP31).*

INTRODUCTION

The Complex I Human Protein Quantity Dipstick Assay Kit (MS131) is used to rapidly quantify the levels of the Complex I enzyme complex in human and bovine sample material. Based on the immunologic sandwich assay, the kit utilizes two monoclonal antibodies (mAbs) specific to different antigens present on the Complex I enzyme complex. One antibody is immobilized on the nitrocellulose membrane in a thin line perpendicular to the length of the dipstick, while the other is gold-conjugated and combined with the sample mix. The sample contents containing the gold-conjugated mAb wick past the mAb immobilized on the dipstick. When assembled Complex I is present in the sample, a red line appears at the site of the anti-Complex I antibody line (Figure 1). The signal intensity is directly related to the amount of Complex I in the sample. The signal intensity is best measured by a dipstick reader (MitoSciences MS1000 Dipstick Reader) or may be analyzed by another imaging system. To identify defects in Complex I that do not affect enzyme assembly combine this assay with MitoSciences' Dipstick Activity Kit for Complex I Activity (MS130) to determine the relative specific activity of Complex I.

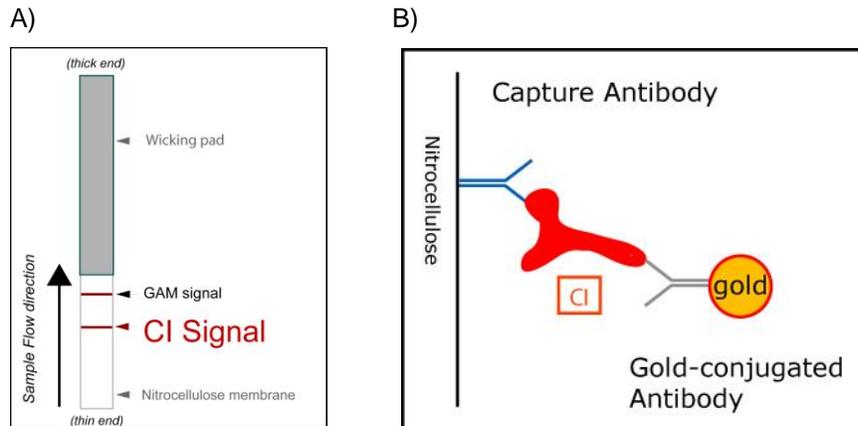


Figure 1. Schematics of a developed Complex I quantity dipstick and the interaction between the mAbs and Complex I. (A) A finished dipstick shows a strong signal for Complex I (bottom band) and the positive control (goat anti-mouse antibody (GAM)) which is used to verify completion of the reaction (i.e. flow of the gold conjugated antibody and sample contents past the capture antibody line). (B) The antibody sandwich diagram showing the interaction between Complex I and the two monoclonal antibodies in this assay.

ADDITIONAL MATERIALS REQUIRED

- Dipstick reader (MitoSciences' MS1000) or other imaging system
- Method for determining protein concentration
- Pipetting devices
- Protease inhibitors

DIPSTICK ASSAY PROTOCOL

A. Sample Preparation

Note: *Samples must be kept on ice. Protease inhibitors are not provided.*

1. Tissue Sample Preparation

- a. Start with approximately 25 mg of tissue sample. Add 10 volumes of Buffer A per microgram of sample (e.g. if the sample weighs 50 mg, add 500 μ L of Buffer A).
- b. Homogenize the sample.
- c. Keep on ice for 20 minutes, mixing intermittently.
- d. Spin the cell extract in a microcentrifuge at 13,000-16,000 rpm at 4°C for 20 minutes to pellet cellular debris.
- e. Transfer supernatant to a new tube and determine protein concentration (e.g. BCA assays are recommended as the detergent in Buffer A does not affect the reading in the assay).
- f. Proceed directly to Part B of the Protocol or freeze samples at -80°C.

2. Cell Culture Sample Preparation

- a. Start with a cell pellet. Add 10 volumes of Buffer A to the cell pellet and mix (e.g. if the total sample volume displaces 50 μ L of volume, then add 500 μ L of Buffer A).
- b. Keep on ice for 20 minutes, mixing intermittently.
- c. Spin the cell extract in a microcentrifuge at 13,000-16,000 rpm at 4°C for 20 minutes to pellet cellular debris.
- d. Transfer supernatant to a new tube and determine protein concentration (e.g. BCA assays are recommended as the detergent in Buffer A does not affect the reading in the assay).
- e. Proceed directly to Part B of the protocol or freeze samples at -80°C.

B. Dipstick Procedure

The assay is most accurate with a user established standard curve for interpolation of the signal intensity. Following the protein concentration ranges as defined in Table 1, generate a standard curve using a positive control sample.

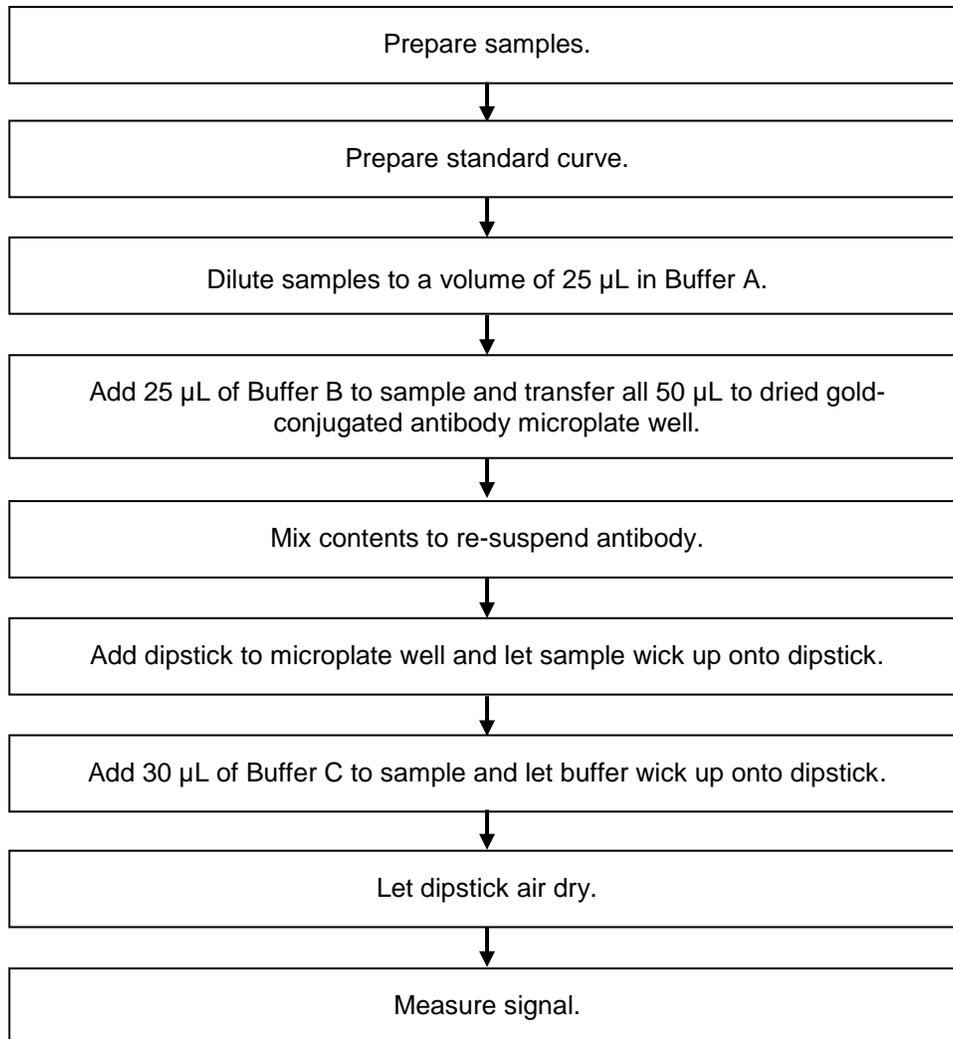
Sample Type	Working Range
Human fibroblasts	0.5 - 20 μ g
PBMC	0.5 - 20 μ g
Lymphoblastoid	0.25 - 10 μ g
Fat	2 - 40 μ g
Bovine heart mitochondria	5 - 300 μ g

Table 1. *Suggested working range for different sample types*

1. Load the amount of protein that corresponds toward the high end of the user generated standard curve (~3/4 of the high end).
2. Add Buffer A to the sample to bring volume up to 25 μ L.
3. Add the sample and 25 μ L of Buffer B to a microplate well of dried gold-conjugated antibody, for a total sample volume of 50 μ L. Incubate the sample for 5 minutes, allowing the gold-conjugate to hydrate.
4. Re-suspend the dried gold-conjugated secondary antibody using a pipette.
5. Gently add a dipstick to the microplate well (place the thin/nitrocellulose end of dipstick down). The dipstick must reach the bottom of the well.
6. Allow the sample wick up onto the dipstick until the entire sample volume has been absorbed (the positive control band typically appears within 2 minutes. A 50 μ L sample should be wicked up completely in 20 minutes; more viscous extracts may take longer. **The dipstick must absorb the entire sample volume before Buffer C is added.**
7. Add 30 μ L of Buffer C and let the entire buffer volume wick up onto the dipstick.
8. Leave dipstick to air-dry (requires approximately 20 minutes to air dry, or 10 minutes in an incubator at 37°C).
9. Measure the signal intensity with a dipstick reader (MitoSciences' MS1000 Dipstick Reader) or other imaging system, e.g. flat-bed scanner.

FLOW CHART

For quick reference only. Be completely familiar with the previous details of this protocol before performing the assay.

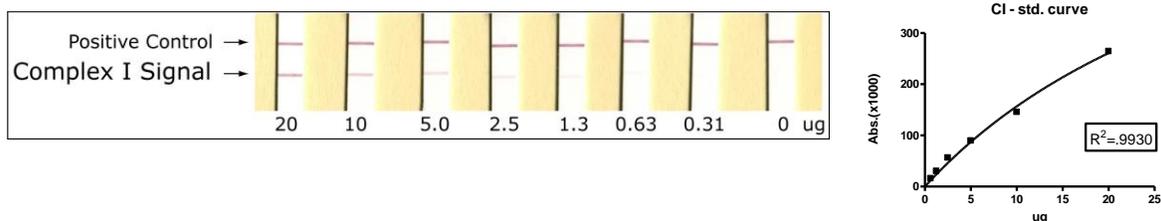


EXAMPLE EXPERIMENT

Illustrated is an experiment using the MS131 assay kit to quantify Complex I levels using various concentration of human fibroblast extract. Samples were prepared as described in this protocol. All data were analyzed using MitoSciences Dipstick Reader (MS1000) and GraphPad software.

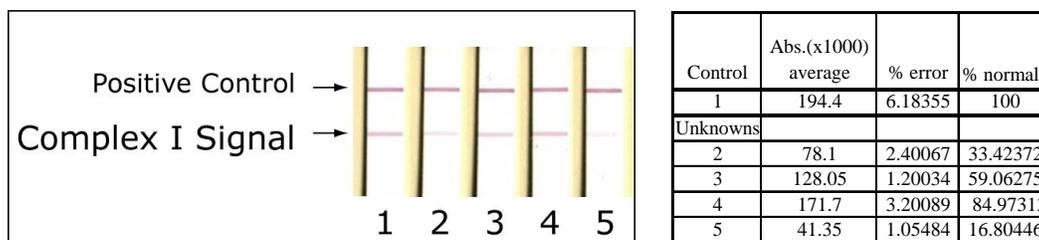
Step 1. Generating a standard curve

Shown is a 1:2 dilution series using a positive control sample (see Table 1 for recommended protein ranges). Approximately 7 to 8 dipsticks are suitable for covering the entire working range and the blank for background levels. In this example the dilution series starts with 20 μg of fibroblast extract. A one-site hyperbola line was generated for best-fit analysis using GraphPad.



Step 2. Analysis of samples

Based on the above standard curve values, 13 μg of protein extract were loaded to each dipstick for the control sample (1) and the four unknown samples (2-5).



Using GraphPad software, the signal intensity from the standard curve was interpolated and the quantity of Complex I in samples 1-5 was determined. Based on the above analysis, the unknown samples had 16% and 85% of Complex I levels compared to the control.

TROUBLESHOOTING GUIDE

Signal is saturated

It is very important that the amount of sample used is within the working range of the assay (a best fit line for interpolation as generated with the GraphPad program). Therefore, it is crucial to know the working range for the sample type and avoid the region of signal saturation (see Table 1). Determination of the working range is recommended for the sample in case of signal saturation.

Signal is too weak

This occurs when the sample lacks measurable amounts of the protein. To increase signal add more sample to another dipstick.

Positive control band does not appear

Make sure that the wicking paper is in contact with the nitrocellulose membrane. Care should always be taken when handling the dipsticks to prevent disruption of this junction (see Figure 1).